0428



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/888,182A

DATE: 05/07/2002 TIME: 14:50:40

Input Set : A:\EP.txt

Output Set: N:\CRF3\05072002\1888182A.raw

## ENTERED

3 <110> APPLICANT: Vertex Pharmaceuticals Incorporated 5 <120> TITLE OF INVENTION: ERK-5 DEFICIENT ANIMALS AND METHODS OF INHIBITING ANGIOGENESIS THROUGH 6 THE INHIBITION OF ERK-5 8 <130> FILE REFERENCE: VPI/00-115 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/888,182A 11 <141> CURRENT FILING DATE: 2001-06-22 13 <150> PRIOR APPLICATION NUMBER: US60/214,044 14 <151> PRIOR FILING DATE: 2000-06-23 16 <160> NUMBER OF SEQ ID NOS: 6 18 <170> SOFTWARE: PatentIn version 3.0 20 <210> SEQ ID NO: 1 21 <211> LENGTH: 26 22 <212> TYPE: DNA 23 <213> ORGANISM: Mus musculus 25 <400> SEQUENCE: 1 26 26 cagccattcg atgtgggccc acgcta 29 <210> SEQ ID NO: 2 30 <211> LENGTH: 25 31 <212> TYPE: DNA 32 <213> ORGANISM: Mus musculus 34 <400> SEQUENCE: 2 25 35 tataacattc tcatggcgga atcgc 38 <210> SEO ID NO: 3 39 <211> LENGTH: 802 40 <212> TYPE: DNA 41 <213> ORGANISM: Mus musculus 43 <220> FEATURE: 44 <221> NAME/KEY: misc\_difference 45 <222> LOCATION: (1)..(802) 46 <223> OTHER INFORMATION: Wherein n is selected from any nucleic acid 49 <400> SEQUENCE: 3 W--> 50 cggnacctac tgtgccctat ggaggaattc agatctgtgt aagggagtgg gccaggagga 60 52 ggagacacag tcgggatcag cttagaagcc caggttcagt aatactgaag ttctggcagg 120 54 geggttgaac ceagagtgat gegggetgtg agteeaggae attggtaggg acagttetta 180 56 teteteaaga gggcaaggge tggggatgte gateaetggt aggetgatga geatetttga 240 300 5/8 ggttttaggt tgactctcct gtacaaaagg ggaaaagaat caagaggatt tacctcttta 60 tggtcatgcc acctttggtt atatcataag ttcaaggcta gtctagaccc tgttccaaaa 360 62 gacaaaacan aaaaccnaaa cagcaatnta nganaaggga gagagggcnc agacngnccg 420 480 64 ggacagatee aaattgtaag acaaeggaca caatacattg tagtgteaca cageagtgte 540 66 ctcatggcag acaactaatt attcacagaa tacctcctta aaaatagagt cttcaacata 600 68 gctttttcag tagctgttgg caaactgtag agtttgctct aaaattaacc atactggcca 660 -> 70 atcttggtag atttgaatat ttctataaaa aaaatttttt ttgacagaaa ttangtccat 720 W--> 72 ggagaaagtg atttgtcaga aagcttgtaa aaaagtttgg ggctnggaaa aaacccgatt

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```
🕉 74 cggtgattaa gatcactcga tcttttaaaa gggacttggc tttaantncc ataatggnct
                                                                               780
                                                                               802
   > 76 ttcaccgggg ggcntaaact tt
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     80 <211> LENGTH: 794
     81 <212> TYPE: DNA
     82 <213> ORGANISM: Mus musculus
     84 <220> FEATURE:
     85 <221> NAME/KEY: misc_difference
     86 <222> LOCATION: (1)..(794)
     87 <223> OTHER INFORMATION: Wherein n is selected from any nucleic acid
     90 <400> SEQUENCE: 4
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W--> 91 gattnaagat cccctcgatn tttnaaaagg acttggnttc aagggaanag ngtnttnncg
                                                                               120
W--> 93 ggggnnaact tgaattggga cnccggtgtt gggatcanac tccctctttn ngcctctgta
                                                                               180
W--> 95 nacccagggc acccaagtag tacacatacg ttcaggaaan catacacata cgtttaagaa
W<sub>1</sub>-> 97 aactttataa aagttgtggc cagncggtgg tggcgcatgc ctttaatccc agcactgggg
                                                                               240
                                                                               300
    /99 aggcagaggc aggcagatct cttgagtttt gggtttgagg ccaacctggt ctacaagagc
                                                                                360
     101 aaqcaaqttc caqqccaqat aaqqctacac agacatcttg tcttgaaaaa aagaaagaaa
                                                                                420
W--> 103 qaatqaaagt tgtagaaaac ctaaaacccg gtgnnnaant ccncncttcc catgntgtta
     105 gtcctttggg gtttcactgt aaggccataa cctcaggaat tgggagtgcc aggggacgga
                                                                                480
     107 gtgccagggg gggcttctcc ctgtgatgtg aggaggctag ctcacccgtt tcttcccatt
                                                                                540
     109 ttcagctatg tggtactgga cctcatggag agcgacctac accagatcat tcactcttca.
                                                                                600
     111 caqccqctca ccctqqaaca tgtgagatac ttcctgtacc agctgcttcg gggcctcaaa
                                                                                660
                                                                                720
     113 tacatgcact ctgctcaggt catccaccgt gatcttaaac cctctaacct tctggtcaat
                                                                                780
     115 gagaactqtq agctcaaqat cqqtqacttt ggaatggccc gtggcctctg tacttccccc
                                                                                794
     117 tgccgagcac caga
     120 <210> SEQ ID NO: 5
     121 <211> LENGTH: 632
     122 <212> TYPE: DNA
     123 <213> ORGANISM: Mus musculus
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     126 <221> NAME/KEY: misc_difference
     127 <222> LOCATION: (1)..(632)
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     131 <400> SEQUENCE: 5
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                                                                                 60
                                                                                120
     134 cacaqttcca cttqaqccca qgqatqqcat ctcaacatct ggacacacag gctcactagc
                                                                                180
W--> 136 cacaggetge ananaagntg gaacgnattg ttgncgaatg ccctccngtc gtgcatgaaa
                                                                                240
     138 gtcttcattc tcaqccacaa tggcctcctt aatgcgctcc ctggtaaggg cttcacgggt
                                                                                300
     140 caaaagcaaa gtcaaaaggt ggggcgcaat caggctcatc atcagggtca tggtacttag
                                                                               360
W--> 142 ccagaagggg tgcgaaggca gcagcagtna gattcgggcn ctgggttcaa ntcgcaccat
     144 gcgtcccagc agggagaggg cctggcggtc agcacctggg tatactgtct cccaaggcac
                                                                                420
     146 aggttgcctt ggtggcaggc tctggatata ggctcgcacc ctttcagccc ccacagcctg
                                                                                480
                                                                                540
     148 aatcacaget ggtgacggag tteecaacae cateatgate agetgtaaet ggtgeaegta
     150 gtttttgcct gggaagaget ggcgccgage cagcatetea ccaaagatge ageccacaga
                                                                                600
                                                                                632
     152 ccagaggtcg attgcctgcg gtatactcgt gc
     155 <210> SEQ ID NO: 6
     156 <211> LENGTH: 617
     157 <212> TYPE: DNA
     158 <213> ORGANISM: Mus musculus
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	160	<220> FEATU	JRE:			•	•	
	161	<221> NAME/	/KEY: misc_d	difference				
	162	<222> LOCATION: (1)(617)						
	163	<223> OTHER	R INFORMATIO	N: Wherein	n is select	ed from any	nucleic acid	
	166	<400> SEQUE	ENCE: 6					
w>	167	ggcaggtacc	gcgttagnac	cnnttatcng	aacconntgt	ttntcncagn	nnagcnntat	60
W>	169	ttaaccttgn	aaanagtttt	tccctgaggc	caagatagca	natangctcn	nnggagnncn	120
			tgttctaaga					180
WF7b(	<u>1</u> 73	antctggaat	gacctnanta	tggtgagtag	gaagggaaga	aggatcagtt	aatncagtca	240
W 17,	175	caancnnntg	ctaactaacg	ngcctcctnt	ttatgtaagc	nattagcanc	ngtttcnnga	300
			aattaaaatn					360
W>	179	acttnatgnn	ntntgggntc	cagaagagca	gtgctgaagg	gacctgcagc	taacttgaag	420
	181	gtactctctg	gtatatgccc	ttttcctgct	ccccaggcca	gcaggtggcc	atcaagaaga	480
	183	tacctaatgc	ttttgatgtg	gtgaccaatg	ccaaacggac	cctcagggag	ctgaagatcc	540
	185	tcaaacactt	caaacacgac	aatatcatcg	ccatcaagga	catcctgaag	cctactgtgc	600
	187	cctatggaga	attcttc					617

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/888,182A

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## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the  $\langle 220 \rangle$  to  $\langle 223 \rangle$  fields of each sequence which presents at least one n or Xaa.

Seq#:3; N Pos. 4,370,377,388,391,394,409,415,417,654,705,766,768,778,794 Seq#:4; N Pos. 5,20,24,37,48,51,54,57,58,65,66,82,98,110,111,121,160,204

Seq#:4; N Pos. 394,395,396,399,403,405,415

Seq#:5; N Pos. 12,14,17,23,42,51,132,134,138,146,154,167,329,340,351

Seq#:6; N Pos. 18,22,23,29,36,37,43,46,50,51,52,56,57,70,74,101,105,110,111
Seq#:6; N Pos. 112,117,118,120,144,145,150,166,170,182,196,198,234,244,246
Seq#:6; N Pos. 247,248,261,269,281,289,291,297,298,320,345,365,369,370,371

Seq#:6; N Pos. 373,378